ENTERED   PCT10

RAW SEQUENCE LISTING   DATE: 03/01/2002
PATENT APPLICATION: US/10/049,868   TIME: 15:02:30

        Input Set : A:\PCT EPO 07874 Sequence listing.txt
        Output Set: N:\CRF3\03012002\J049868.raw

```
   3 <110> APPLICANT: K. U. Leuven Research & Development
   4         Deckmyn, Hans
   5         Cauwenberghs, Nancy
   7 <120> TITLE OF INVENTION: Cell lines, ligands and antibody fragments for use in
   8         pharmaceutical compositions for preventing and treating
   9         haemostasis disorders
  11 <130> FILE REFERENCE: 6B4FAB
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/049,868
C--> 13 <141> CURRENT FILING DATE: 2002-02-11
  13 <150> PRIOR APPLICATION NUMBER: EP00102032.0
  14 <151> PRIOR FILING DATE: 2000-02-02
  16 <150> PRIOR APPLICATION NUMBER: GB9918788.2
  17 <151> PRIOR FILING DATE: 1999-12-10
  19 <160> NUMBER OF SEQ ID NOS: 4
  21 <170> SOFTWARE: PatentIn version 3.1
  23 <210> SEQ ID NO: 1
  24 <211> LENGTH: 359
  25 <212> TYPE: DNA
  26 <213> ORGANISM: Mus sp.
  28 <220> FEATURE:
  29 <221> NAME/KEY: V_region
  30 <222> LOCATION: (1)..(359)
  31 <223> OTHER INFORMATION:
  33 <220> FEATURE:
  34 <221> NAME/KEY: primer_bind
  35 <222> LOCATION: (1)..(24)
  36 <223> OTHER INFORMATION: Vkback primer
  38 <220> FEATURE:
  39 <221> NAME/KEY: primer_bind
  40 <222> LOCATION: (328)..(359)
  41 <223> OTHER INFORMATION: Vk2for primer
  43 <220> FEATURE:
  44 <221> NAME/KEY: misc_feature
  45 <222> LOCATION: (70)..(100)
  46 <223> OTHER INFORMATION: complementary determining region one
  48 <220> FEATURE:
  49 <221> NAME/KEY: misc_feature
  50 <222> LOCATION: (144)..(165)
  51 <223> OTHER INFORMATION: complementary determining region two
  53 <220> FEATURE:
  54 <221> NAME/KEY: misc_feature
  55 <222> LOCATION: (262)..(289)
  56 <223> OTHER INFORMATION: complementary determining region three
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 58 <400> SEQUENCE: 1
 59 gacattgagc tcacccagtc tccagcaatc atgtctgcat ctccagggga gaaggtcacc      60
 61 atgacctgca gtgccagttc aagtgttaat tacatgcact ggttccagca ggagtcgggc     120
 63 accttcccca aaagaaggat ttatgacaca tccaaactgg cttctggagt ccctgctcgc     180
 65 ctcagtggca gtgggtctgg gacagaattc accctggaaa tcagtagagt gaaggctgag     240
 67 gatgtgggtg tgtattactg tcaacaactt gtagagtatc cgctcacgtt cggtgctggg     300
 69 accaagctgg agctgaaacg ggctgatgct gcaccaactg tatccatctt caagcttcc      359
 72 <210> SEQ ID NO: 2
 73 <211> LENGTH: 363
 74 <212> TYPE: DNA
 75 <213> ORGANISM: Mus sp.
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W--> 76 **<220> FEATURE:**
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 77 <221> NAME/KEY: V_region
 78 <222> LOCATION: (1)..(363)
 79 <223> OTHER INFORMATION:
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 82 <221> NAME/KEY: primer_bind
 83 <222> LOCATION: (1)..(24)
 84 <223> OTHER INFORMATION: Vhback primer
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W--> 86 **<220> FEATURE:**
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 87 <221> NAME/KEY: primer_bind
 88 <222> LOCATION: (330)..(360)
 89 <223> OTHER INFORMATION: Vhfor primer
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W--> 91 **<220> FEATURE:**
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 92 <221> NAME/KEY: misc_feature
 93 <222> LOCATION: (90)..(105)
 94 <223> OTHER INFORMATION: complementary determining region one
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W--> 96 **<220> FEATURE:**
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 97 <221> NAME/KEY: misc_feature
 98 <222> LOCATION: (147)..(195)
 99 <223> OTHER INFORMATION: complementary determining region two
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W--> 101 **<220> FEATURE:**
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102 <221> NAME/KEY: misc_feature
103 <222> LOCATION: (291)..(330)
104 <223> OTHER INFORMATION: complementary determining region three
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109 acttgcactg tctctgggat ttcattaaac agatatggtg tacactgggt tcgccagcct     120
111 ccaggaaagg gtctggagtg gctgggagta atatggactg gtggaagcac aaattataat     180
113 tcggctctca tgtccagact gagcatcagc aaagacaact ccaagagcca agtttttctta    240
115 aaaatgaaca gtctgcagac tgatgacaca gccatgtact actgtgccag agatcgatct     300
117 actatgatta cggcctatgc tatggactac tggggccaag gaccacggt caccgtctcc      360
119 tca                                                                   363
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123 <211> LENGTH: 119
124 <212> TYPE: PRT
125 <213> ORGANISM: Mus sp.
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W--> 126 **<400> SEQUENCE: 3**
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128 Asp Ile Glu Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Pro Gly
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```
129 1                    5                    10                   15
132 Glu Lys Val Thr Met Thr Cys Ser Ala Ser Ser Ser Val Asn Tyr Met
133              20                  25                  30
136 His Trp Phe Gln Gln Glu Ser Gly Thr Phe Pro Lys Arg Arg Ile Tyr
137              35                  40                  45
140 Asp Thr Ser Lys Leu Ala Ser Gly Val Pro Ala Arg Leu Ser Gly Ser
141          50                  55                  60
144 Gly Ser Gly Thr Glu Phe Thr Leu Glu Ile Ser Arg Val Lys Ala Glu
145 65                  70                  75                  80
148 Asp Val Gly Val Tyr Tyr Cys Gln Gln Leu Val Glu Tyr Pro Leu Thr
149              85                  90                  95
152 Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys Arg Ala Asp Ala Ala Pro
153              100                 105                 110
156 Thr Val Ser Ile Phe Lys Leu
157          115
160 <210> SEQ ID NO: 4
161 <211> LENGTH: 121
162 <212> TYPE: PRT
163 <213> ORGANISM: Mus sp.
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W--> `164 <400> SEQUENCE: 4`
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167 1                    5                    10                   15
170 Ser Leu Ser Ile Thr Cys Thr Val Ser Gly Ile Ser Leu Asn Arg Tyr
171              20                  25                  30
174 Gly Val His Val Arg Gln Pro Pro Gly Lys Gly Leu Glu Trp Leu
175              35                  40                  45
178 Gly Val Ile Trp Thr Gly Gly Ser Thr Asn Tyr Asn Ser Ala Leu Met
179          50                  55                  60
182 Ser Arg Leu Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln Val Phe Leu
183 65                  70                  75                  80
186 Lys Met Asn Ser Leu Gln Thr Asp Asp Thr Ala Met Tyr Tyr Cys Ala
187              85                  90                  95
190 Arg Asp Arg Ser Thr Met Ile Thr Ala Tyr Ala Met Asp Tyr Trp Gly
191              100                 105                 110
194 Gln Gly Thr Thr Val Thr Val Ser Ser
195          115                 120
```

VERIFICATION SUMMARY DATE: 03/01/2002
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```
L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:76 M:283 W: Missing Blank Line separator, <220> field identifier
L:126 M:283 W: Missing Blank Line separator, <400> field identifier
L:164 M:283 W: Missing Blank Line separator, <400> field identifier
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